

## **RAW SEQUENCE LISTING**

**The Biotechnology Systems Branch of the Scientific and Technical Information Center (STIC) no errors detected.**

Application Serial Number: 10/535, 523 A  
Source: PCT  
Date Processed by STIC: 09/26/2005

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RAW SEQUENCE LISTING  
PATENT APPLICATION: US/10/535,523A

DATE: 09/26/2005  
TIME: 12:50:39

Input Set : D:\seq listing.app  
Output Set: N:\CRF4\09262005\J535523A.raw

3 <110> APPLICANT: The Government of the United States of America  
5 <120> TITLE OF INVENTION: CELL LINES AND HOST NUCLEIC ACIDS RELATED TO INFECTIOUS  
DISEASES

7 <130> FILE REFERENCE: 6395-66741-06  
C--> 9 <140> CURRENT APPLICATION NUMBER: US/10/535,523A  
C--> 9 <141> CURRENT FILING DATE: 2005-05-17  
9 <150> PRIOR APPLICATION NUMBER: PCT/03/037143  
10 <151> PRIOR FILING DATE: 2003-11-18  
12 <150> PRIOR APPLICATION NUMBER: US 60/482,604  
13 <151> PRIOR FILING DATE: 2003-06-25  
15 <150> PRIOR APPLICATION NUMBER: US 60/427,464  
16 <151> PRIOR FILING DATE: 2002-11-18  
18 <160> NUMBER OF SEQ ID NOS: 845  
20 <170> SOFTWARE: PatentIn version 3.2  
22 <210> SEQ ID NO: 1  
23 <211> LENGTH: 937  
24 <212> TYPE: DNA  
25 <213> ORGANISM: Homo sapiens  
28 <220> FEATURE:  
29 <221> NAME/KEY: misc\_feature  
30 <222> LOCATION: (1)..(937)  
31 <223> OTHER INFORMATION: n is a, g, c, or t  
33 <400> SEQUENCE: 1

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35 cgaaaacgatc ctyrgstacc gcttgccaaa cctacaggtg gggctttca agaggcttcc 120  
36 agaccttaggg gagcatctca gcgtcactcg ctgtccagtt gctgtgatca ggtgctttgg 180  
37 ggtttgtgtg actccagaat ccactgggcc ttgtgttcag aagacaaaag ttaaccataa 240  
38 ggcacagaag aaagcctctt gctgaagcca tcgttggccc acatgcattt caggacaag 300  
39 aaatgaagat cggagactt caagttgtgc ccaggactca cctgctccca ggagacaaaa 360  
40 ggcacacacag cagaggagcc tgaagcccat ggcaggatct cctagcttgg ggctgggtgtc 420  
41 tctgttagtaa gcattctgaa gttcctaagg tcccttctt cttgataggag cattgacactg 480  
42 ttagtgtcacc acactgacat actttccctt gcaggccact ccagccact gtactctttg 540  
43 gcaggcctca ggttctgtca ctccatgtac tattctgtc ttgcacaggg cagaagctaa 600  
44 aggtgaggag gactgaacac agtaccaaca taccacatc acaccttact ttccctctgcc 660  
45 cgcctgtcc ctgcctgtac actgattccc cagcccttgc caccacccggc ctttacccct 720  
46 ccactgcccgg tgcagcagca gagacactcc ctccttgatg caaactgagg cctctggcac 780  
47 cccaaacttt tcaaggcaat gatagtctgt gcttaactct acatggccag gcccccaactc 840  
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49 accccgaaga aaagaancaa nngatctttt ggnnacc 937  
50 <210> SEQ ID NO: 2  
51 <211> LENGTH: 1515  
52 <212> TYPE: DNA  
53 <213> ORGANISM: Homo sapiens  
54 <220> FEATURE:

(pg 617)

RAW SEQUENCE LISTING  
PATENT APPLICATION: US/10/535,523A

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74 <221> NAME/KEY: misc\_feature  
75 <222> LOCATION: (1)..(1515)  
76 <223> OTHER INFORMATION: n is a, c, g, or t  
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83 tatgcagtgc tgccataacc atgagtgata acactgcggc caacttactt ctgacaacga 180  
85 tcggaggacc gaaggagcta accgctttt tgcacaacat gggggatcat gtaactcgcc 240  
87 ttgatcggtt ggaaccggag ctgaatgaag ccataccaaa cgacgagcgt gacaccacga 300  
89 tgcctgcage aatggcaaca acgttgcgc aactattaac tggcgaacta cttactctag 360  
91 cttccggca acaattaata gactggatgg aggcgataa agttgcagga ccacttctgc 420  
93 gctcgccct tccggctggc tggtttattt ctgataaattc tggagccggt gagcgtgggt 480  
95 ctcgcggat cattgcagca ctggggccag atggtaagcc ctcccgatc gttagttatct 540  
97 acaacgacggg gagtcaaggca actatggatg aacgaaatag acagatcgct gagataggtg 600  
99 cctcaactgat taagcatting gkaanctgtc agaccattkt ttactgcata tacsgatcca 660  
101 ttgccttat ctcaaactct tattatgaaa tcactnccct tgagagaraa aaagcctttt 720  
103 tctctnnngg atkgtcccag magytyccga mcatccccac tycccaacct tatgkggccc 780  
105 agcaatgans cctagtagta ggaaaatcty tatggatacy ggkgnctgak ggaaaratcc 840  
107 ttcytctcat gaarwgtatgg kgactggggc tytggatgc tcacggaaat ccctatttcc 900  
109 cccacaaaaga agttattttta ttacacaacc atttggatga cccccctttt ctccaattt 960  
111 nccaaataaaa tctgtaaagg tcacaggtga agttcttctc tttaaagagct actccatgct 1020  
113 aagttcagcg agaacttggg gtaccctaga cattttccca gagatgcctt tcttgcact 1080  
115 ctttcaata agtaagcatg ctttgcctg cactgggtgt cacctgtt ggatgctgtt 1140  
117 gtcctgcct tgccttat tctgtccaca tggtttctc ataggatgt gcttaggtca 1200  
119 gcccggat ttgaaccagt caacaagtcc aggttgggtgt ggagccctt tagtacctcc 1260  
121 ctttgcagga ataatgtgc acccagaaac tccctcagag cctctccact gggagggggcc 1320  
123 ttgtgaccat tcctggtttta ctcccttctgt tccagcatcc catgtggcca atggggccct 1380  
125 ttcattttca atggatctc antnttaca gtaagttata ttattgcct acatngaact 1440  
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129 anaatccact tcgnn 1515

132 <210> SEQ ID NO: 3  
133 <211> LENGTH: 885  
134 <212> TYPE: DNA  
135 <213> ORGANISM: Homo sapiens  
138 <220> FEATURE:  
139 <221> NAME/KEY: misc\_feature  
140 <222> LOCATION: (1)..(885)  
141 <223> OTHER INFORMATION: n is a, g, c, or t  
143 <400> SEQUENCE: 3

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146 cgagttcttc tgagcgggac tctgggggtc gaaatgagct agcccttaag taacgccatt 120  
148 ttgcaaggca tggaaaata cataactgag aatagaaaag ttcagatcga ggtcaggaac 180  
150 agatggaaaca gggtcgaccg gtcgaccggt cgaccctaga gaaccatcag atgttccag 240  
152 ggtcccccaa ggacctgaaa tgaccctgtg ctttatttga actaaccaat cagttcgctt 300  
154 ctcgcctctg ttcgcgcgt tctgcctccc gagctcaata aaagagccca caaccctca 360  
156 ctcggggcgc cagtctccg attgactgag tgcggccgggt acccggtat ccaataaacc 420  
158 ctcttgcagt tgcattccgac ttgtggtctc gctttccctt gggaggggtct cctctgagtg 480  
160 attgactacc cgtcagcggg ggtcttccac tctctgtgtt ctggatccaa cagagcctgg 540  
162 accagggccct ccagttccctc attcgttattt ataatggaga agagagagca aaaggaaaca 600

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164 ttcttgaacg attctccgca caacagttcc ctgacttgca ctctgaacta aacctgagct	660
166 ctctggagct gggggactca gctttgtatt tctgtgccag cagcgttaggt ggttagcttga	720
168 aacagttctt cgggcccagg acacggctca ccgtgctagg taagaagggg gctccagtgg	780
170 gagagagggg gagcagccca ncctgnncga ccccanancc tggtnnttagg ggagtggnc	840
172 ctgggcatcc aggcctnct cnaggaancg ggttncgccc ggncc	885
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183 <222> LOCATION: (1)..(900)	
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189 acatagaaaa aaatctgaaa agatatctca aaagccaga catttatca cactaacgg	120
191 gaaaagcata ccccacagtg tcagtggagg caacatgggg tcctggattt ccttccacc	180
193 ctcagtggta gtgaggtgtt cctctcactc cttctgagta gaggaagcca agaggaaagc	240
195 tggaaacttgt accatcatcc agtggtgata aagcctctgt ccctccacct taccccccagg	300
197 ttatcagtgg caaccacatg gctagtggta cccctcccgcc tcctagccag aatgatata	360
199 gcagaggcct agagagtagc ccaaaaactc atctgcaccc agcaggactg aggtttccta	420
201 ccccccaccaa tggaaagccaa gtgaggaacc taaggcattca cctctcactc agcaggaacc	480
203 agacaacacc ccctaacaaca cacacacaca cacacacaca cacacccttc tgtagtgt	540
205 gtatcaagga ggcttgataa aatagaagat ttaaatagga tccattgccc ttatctcaaa	600
207 ctcttattat gaaatcactc ctttgagaga gaaaaaaagcc ttttcttggc ggattgtccc	660
209 agcagctccc gaccatcccc actcccaac cttatgtggc cccagcaatg agcttagtag	720
W--> 211 taggaaaatc tctatggata ctgggtctga tggaaagatt cttcctctca ngaagtgtat	780
213 gtgactgggg ctctggatg ctcacgggaa tnccatttcc cccacaagaa ntattttat	840
215 naccaaccat ttggatgacc ctttttntt ccatttncca annaatttgt aagncaag	900
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219 <211> LENGTH: 869	
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221 <213> ORGANISM: Homo sapiens	
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232 ttcttccaaat ttcccaaata aatctgtaaa ggtcacaggt gaagttcttc tcttaagag	120
234 ctaactccatg ctaagttcag cgagaacttg gggtaaccctt gacatttttc cagagatgt	180
236 ttcttgcata ctctttcaa taagtaagca tgcttgctc tgcaactgggt gtcacccgt	240
238 ttggatgctt ttgtccctgc cttggccctt attctgttca catggtttct tcataaggat	300
240 atgcttaggt cagccctgag gtttgaacca gtcaacaatg ccagggttggt gtggagttccc	360
242 tttagtaccc cccttgcag gaataatgt gcacccagaa actcccttag agcctctcca	420
244 ctggaggggc cttgtgacca ttcctgggtt actcccttgc ttccagcatc ccatgtggcc	480
246 aatggggcccc ttccattttc aatggtatct caattttac agtaagttat attattgtccc	540
248 tacatcgaac tcatttttc tcagtgttac ctgaggaaga atggagagga tgcccagaat	600
250 tggcccagaa gaatccactt cgattctaga gaaaaaggca ggttagaggca gaagagattc	660

RAW SEQUENCE LISTING DATE: 09/26/2005  
 PATENT APPLICATION: US/10/535,523A TIME: 12:50:39

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254	ctgtaaaact	tggaaagagg	aacctgccct	gggtcaagta	gggtgttggg	aggaccagat	780
256	ggagcttcaa	gctctctcca	tctttgtcaa	gtccctgga	ctgagagggn	aaaatnacat	840
258	ggcctttatac	ctccagagga	aantnattc				869
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268	<221>	NAME/KEY:	misc_feature				
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270	<223>	OTHER INFORMATION:	n is a, g, c, or t				
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277	tgcaaggcat	ggaaaaataac	ataactgaga	atagaaaaat	tcaagatcgag	gtcaggaaca	180
279	gatggaacag	ggtcgaccgg	tcgaccggtc	gaccctagag	aaccatcaga	tgttccagg	240
281	gtgccccaaag	gacctgaaaat	gaccctgtgc	cttatttcaa	ctaaccatac	agttcgcttc	300
283	tcgcttctgt	tcgcgcgctt	ctgctcccg	agctcaataa	aagagccac	aacccttcac	360
285	tcggggcgcc	agtccctccga	ttgactgagt	cgcccggtt	cccggttatac	caataaaaccc	420
287	tcttgcagtt	gatccgact	tgtggtctcg	ctgttccttgc	ggagggttctc	ctctgagtga	480
289	ttgactaccc	gtcagcgcccc	gtctttcaact	ctctgtgtac	tggtaccaac	agagcctgga	540
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293	tcttgaacga	ttctccgcac	aacagttccc	tgacttgcac	tctgaactaa	acctgagctc	660
295	tctggagctg	ggggactcag	ctttgtattt	ctgtgccagc	agcgttaggt	gtagcttcaa	720
297	acagttcttc	gggcccaggaa	cacggctcac	cgtgttaggt	aagaaggggg	ctccaggtgg	780
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320	actccagaat	ccactggggcc	tgtgtgtca	aagacaaaag	ttaaccataa	ggcacagaag	180
322	aaaggctcc	gctgaagcca	tcgttggccc	acatgcattt	caggacaaag	aatgaagat	240
324	cggagactt	caagttgtgc	ccaggactca	cctgtccca	ggagacaaaa	ggccacacag	300
326	cagaggagcc	tgaagccat	ggcaggatct	cctagcttgg	ggctgggttc	tctgttagtaa	360
328	gcattctgaa	gttcctaagc	tccctcttc	ctgataggag	cattgacact	tgtatgtcacc	420
330	acactgacat	actttccct	gcaggccact	ccagccact	gtactcttgc	gcaggcctca	480
332	ggttctgcta	ctccatgtac	tattcctgtc	ttgcacaggc	cagaagctaa	aggtgaggag	540
334	gactgaacac	agtaccaaca	tacccacatc	acacctact	ttcctctgccc	cgcctgtcc	600
336	ctgcccctgac	actgattccc	cagcccttgc	cacccagcc	ccttcaccc	ccactgccc	660
338	tgcagcagca	gagacactcc	ctccttgatg	caaactgagg	cctctggcac	cccaactctt	720

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340	tcagggcaat gatagtctgt gcttaactct acatggccag gccccactca gggattctc	780
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344	tacagac	847
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355 <222>	LOCATION: (1)..(755)	
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361	taagtgcgaaat ttatttttt tttttttt ttttttggaa cagagtctcg ctctgtcacc	120
363	caggcttagag tgcagtggca tgatcttggc tcactgcaac ctccacccctc cagttcaag	180
365	tgattctctt gcctcagcct cccaagttagc tagtattaca gacgcctgac accacgccccg	240
367	gttaatttttt gtacttttag tagagacagg tttcaccata ttggccagggc tggctctaaa	300
369	ctcctgaccc caggtgatcc tcctgcctca gcctccaaa gtgctggat tacaggcatg	360
371	agctaccacg tctggctaa gtgcgttta cctatactaa caaaaaccaca cttctgcctc	420
373	gaatgagaac agtctcttgc acatcttggc tctttgcctg actcaaaagcc tcaggtctaa	480
375	gcctcccccatt aatttcttagt ctcagcagaa agatcaatga caggagactc tccaggtat	540
377	gaaaattaaacc aattaagtaa cttgggttgg catctcccg tttgttccacc agtcacactn	600
379	ctgccacagg tatatccctt ctctcancca tatatgcaca aaccccccnc ccacggnaca	660
381	catannaana atttggaaaga ctanaaaatc aggcaangtn tanncncacct tgnnggctgg	720
383	agtatggnan cttggccgg nacatncata cattg	755
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400	accttcaacta caaggatatt acaacaggac attttttaaa acctcaaaaca tcacccaaat	120
402	ttctaagtgc aagtttattt ttatttttt tttttttt gagacagagt ctcgtctgt	180
404	cacccaggct agagtgcagt ggcgtatct tggctcactg caacccctc ac ctcagggt	240
406	caagtgattt tcttgcctca gcctcccaag tagcttagat tacagacgcc tgccaccacg	300
408	cccggttaat ttttgtactt ttagtagaga caggttccat catattggcc aggtggct	360
410	caaactcctg acctcagggtg atcctcctgc ctcaggctcc caaagtgcgt ggattacagg	420
412	catgagctac cacgtctggc ctaagtgcattt gttacctata ctaacaaaac cacacttctg	480
414	cctcgaatga gaacagtctc ctgaacatct tgcctctttg cctgactcaa agcctcagg	540
416	ctaaggctcc ccataatttc tagtctcagc agaaagatca atgacaggag actctccagg	600
418	tgtatggaaattt aaccaattaa gtaacctggg ttggcatctt cccgtttttt caccagctca	660
420	cctcctgcca caggtatatac ctttctctca gccatatacg cacaaccccc ctcacccacgg	720
422	cacacataga aanaatttgg aagactagaa aatcaggcn a gggnttanca caccctngag	780
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**RAW SEQUENCE LISTING ERROR SUMMARY**      **DATE:** 09/26/2005  
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**Please Note:**

Use of n and/or Xaa have been detected in the Sequence Listing. Please review the Sequence Listing to ensure that a corresponding explanation is presented in the <220> to <223> fields of each sequence which presents at least one n or Xaa.

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Seq#:3; N Pos. 9,801,806,807,816,818,825,838,858,862,868,875,880,883  
Seq#:4; N Pos. 771,812,831,841,868,877,882,883,895  
Seq#:5; N Pos. 6,8,21,830,836,863,865  
Seq#:6; N Pos. 801,816,836,847  
Seq#:7; N Pos. 797  
Seq#:8; N Pos. 7,15,600,627,649,657,665,666,669,684,696,700,703,705,713,728  
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Seq#:10; N Pos. 21,584,624,670,696,698,711,727,745,756,768,782,795,801,804  
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Seq#:11; N Pos. 9,10,26,39,684,689,694,697,700,703  
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Seq#:13; N Pos. 740  
Seq#:14; N Pos. 11,27,40,700,706,716,745,749,761,790  
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Seq#:15; N Pos. 705,713,723,726,734,736,744,745,746,749,755,762,763,767,777  
Seq#:15; N Pos. 782,784  
Seq#:16; N Pos. 11,570,638,642,654,664,682,689,697,704,711,714,723,742,743  
Seq#:16; N Pos. 754  
Seq#:17; N Pos. 2,3,22,86,695,741  
Seq#:18; N Pos. 7,8,636,693,710,736,748,750,756,762,764,769  
Seq#:19; N Pos. 7,9,521,568,621,629,632,637,641,648,653,659,663,672,687,695  
Seq#:19; N Pos. 709,712,715,721,733,739,745,747,758,759,763,764,766,768  
Seq#:20; N Pos. 6,8,733,746,752,759,760,790,803,807,821,822,823,828,830,847  
Seq#:20; N Pos. 848,856,860,861,869,872,874,876,881,882,895,912  
Seq#:21; N Pos. 2,7,36,723,724,778,799,817,1461,1553,1586,1597,1599,1604  
Seq#:22; N Pos. 8,22,712,798,837  
Seq#:23; N Pos. 8,13,14,22,44,47,54,73,76,87,117,129,1519,1534,1539,1547  
Seq#:23; N Pos. 1548,1550,1558,1560  
Seq#:24; N Pos. 498,503,506,652,655,656,658,794,821,834,858,1335,1368,1375  
Seq#:24; N Pos. 1402,1435,1437  
Seq#:25; N Pos. 8,9,15,32,680,683,700,732,736,745,754,763,764,774,787,797  
Seq#:25; N Pos. 800,805,809,814,815,816,819,830  
Seq#:26; N Pos. 5,740,744,746,770,778,784,785,795,802,820,827,839,842,845  
Seq#:26; N Pos. 852,854  
Seq#:27; N Pos. 6,7,234,235,236,797,807,822,830,839,845,864,870,873  
Seq#:28; N Pos. 3,9,13,16,17,18,19,20,21,24,27,30,31,32,33,34,39,43,47,49  
Seq#:28; N Pos. 50,51,54,56,57,798,806,820,833,853,855,864,865,878,880,884

RAW SEQUENCE LISTING ERROR SUMMARY DATE: 09/26/2005  
PATENT APPLICATION: US/10/535,523A TIME: 12:50:40

Input Set : D:\seq\_listing.app  
Output Set: N:\CRF4\09262005\J535523A.raw

Seq#:28; N Pos. 885,891,894,900,901  
Seq#:29; N Pos. 2,8,9,18,23,26,35,41,50,809,813,826  
Seq#:30; N Pos. 2,3,4,14,18,19,20,27,28,29,31,32,34,38,41,48,49,50,65,72,85  
Seq#:30; N Pos. 90,99,771,782,789,822,827,848,858,862,872,873,879,881,884  
Seq#:30; N Pos. 885,888,889  
Seq#:31; N Pos. 12,21,574,604,688,691,713,715  
Seq#:32; N Pos. 6,250

### Invalid <213> Response:

Use of "Artificial" only as "<213> Organism" response is incomplete, per 1.823(b) of New Sequence Rules. Valid response is Artificial Sequence.

Seq#:238,239,240,241

## VERIFICATION SUMMARY

PATENT APPLICATION: US/10/535,523A

DATE: 09/26/2005

TIME: 12:50:40

Input Set : D:\seq listing.app

Output Set: N:\CRF4\09262005\J535523A.raw

L:9 M:270 C: Current Application Number differs, Replaced Current Application No  
L:9 M:271 C: Current Filing Date differs, Replaced Current Filing Date  
L:34 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:1 after pos.:0  
M:341 Repeated in SeqNo=1  
L:79 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:2 after pos.:0  
M:341 Repeated in SeqNo=2  
L:144 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:3 after pos.:0  
M:341 Repeated in SeqNo=3  
L:211 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:4 after pos.:720  
M:341 Repeated in SeqNo=4  
L:230 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:5 after pos.:0  
M:341 Repeated in SeqNo=5  
L:299 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:6 after pos.:780  
M:341 Repeated in SeqNo=6  
L:342 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:7 after pos.:780  
L:359 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:8 after pos.:0  
M:341 Repeated in SeqNo=8  
L:398 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:9 after pos.:0  
M:341 Repeated in SeqNo=9  
L:439 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:10 after pos.:0  
M:341 Repeated in SeqNo=10  
L:480 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:11 after pos.:0  
M:341 Repeated in SeqNo=11  
L:517 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:12 after pos.:0  
M:341 Repeated in SeqNo=12  
L:556 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:13 after pos.:0  
M:341 Repeated in SeqNo=13  
L:595 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:14 after pos.:0  
M:341 Repeated in SeqNo=14  
L:636 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:15 after pos.:0  
M:341 Repeated in SeqNo=15  
L:677 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:16 after pos.:0  
M:341 Repeated in SeqNo=16  
L:716 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:17 after pos.:0  
M:341 Repeated in SeqNo=17  
L:757 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:18 after pos.:0  
M:341 Repeated in SeqNo=18  
L:796 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:19 after pos.:0  
M:341 Repeated in SeqNo=19  
L:835 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:20 after pos.:0  
M:341 Repeated in SeqNo=20  
L:880 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:21 after pos.:0  
M:341 Repeated in SeqNo=21  
L:947 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:22 after pos.:0  
M:341 Repeated in SeqNo=22  
L:990 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:23 after pos.:0  
M:341 Repeated in SeqNo=23  
L:1073 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:24 after pos.:480

**VERIFICATION SUMMARY**

PATENT APPLICATION: US/10/535,523A

DATE: 09/26/2005

TIME: 12:50:40

Input Set : D:\seq listing.app

Output Set: N:\CRF4\09262005\J535523A.raw

M:341 Repeated in SeqNo=24

L:1120 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:25 after pos.:0

M:341 Repeated in SeqNo=25

L:1161 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:26 after pos.:0